Enhancing Hit Identification in *Mycobacterium tuberculosis* Drug Discovery Using Dual-Event Bayesian Models

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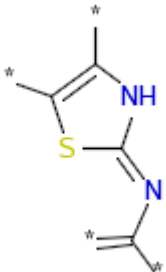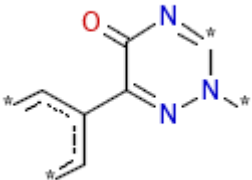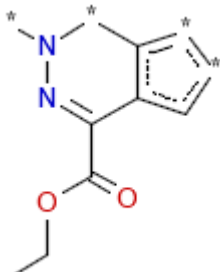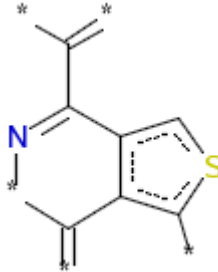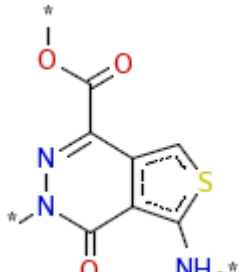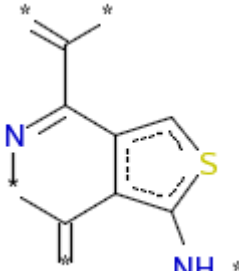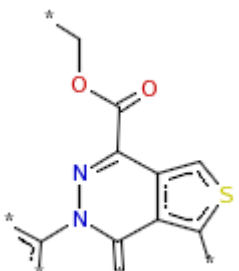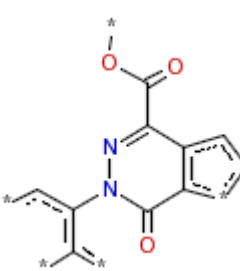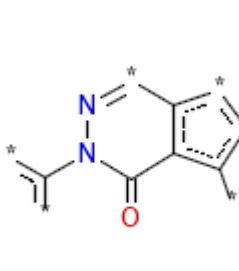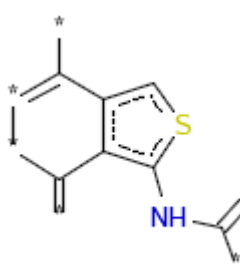
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Running Head: Dual Event Bayesian Models

**Figure S4.** TB kinase single point model: bad features from FCFP_6.

B1: 78996860
0 out of 649 good
Bayesian Score: -3.574

B2: 1780681940
0 out of 609 good
Bayesian Score: -3.512

B3: -186241159
0 out of 608 good
Bayesian Score: -3.510

B4: -163956986
0 out of 608 good
Bayesian Score: -3.510

B5: 1488194790
0 out of 608 good
Bayesian Score: -3.510

B6: 1642535433
0 out of 608 good
Bayesian Score: -3.510

B7: 792169024
0 out of 608 good
Bayesian Score: -3.510

B8: 1040673898
0 out of 608 good
Bayesian Score: -3.510

B9: 389259921
0 out of 608 good
Bayesian Score: -3.510

B10: 229482314
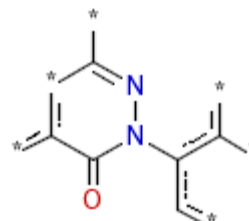0 out of 608 good
Bayesian Score: -3.510
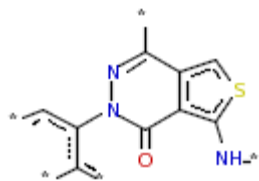
B11: 10587343
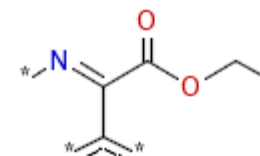0 out of 608 good
Bayesian Score: -3.510
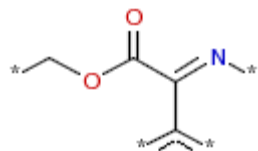
B12: 909560912
0 out of 608 good
Bayesian Score: -3.510
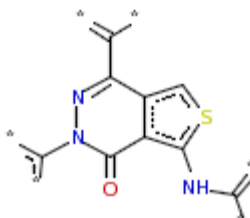
B13: -88401076
0 out of 608 good
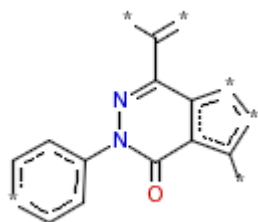Bayesian Score: -3.510

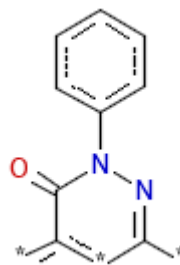B14: -237342290
0 out of 608 good
Bayesian Score: -3.510

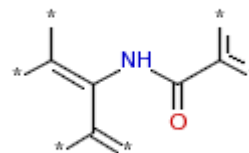B15: 859865998
0 out of 608 good
Bayesian Score: -3.510

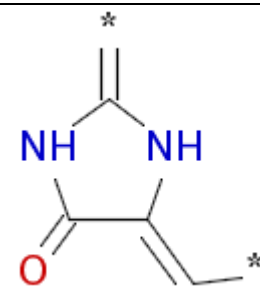B16: -223337671
0 out of 605 good
Bayesian Score: -3.505

B17: -118187393
0 out of 605 good
Bayesian Score: -3.505

B18: 1561760743
0 out of 590 good
Bayesian Score: -3.481

B19: 150315108
0 out of 515 good
Bayesian Score: -3.350

B20: -971446717
0 out of 513 good
Bayesian Score: -3.346